SEQUENCE LISTING

5	Ji: Zho Ji:	un Wang, Benjamin XY Li, Xin Cheng, ng Liu, Li-Wen Niu, Wan-Zhi Huang, en-Yu Xu, Dan Luo, Lian-Di Kang, n-Guo Ding, Fang Rong, Yan Liu, i-Ran Chen
10	(ii) TITLE OF INVENTION	N: AN ANTITHROMBOSIS ENZYME FROM THE SNAKE VENOM OF AGKISTRODON ACUTUS
	(iii) NUMBER OF SEQUENC	CES: 4
15	(B) STREET: 633 (C) CITY: Los	on & Lyon B West Fifth Street B Angeles Lifornia
	(v) COMPUTER READABLE F	ORM:
25 <u>.</u>	(A) MEDIUM TYPE:	3.5" Diskette, 1.44 Mb
•	(B) COMPUTER:	IBM compatible
30	(C) OPERATING SYSTEM:	IBM P.C. DOS (Version 5.0)
	(D) SOFTWARE:	WordPerfect (Version 5.1)
35	<pre>(vi) CURRENT APPLICATION DAT (A) APPLICATION NUMBER (B) FILING DATE: (C) CLASSIFICATION:</pre>	
40	(viii) ATTORNEY/AGENT INFORM (A) NAME: (B) REGISTRATION NUMBE	Chen, Anthony C.
		•

	(ix) TELECOMMUNICATION INFORMA	TION:	
5	(B) TELEFAX:	(213) 489-1600 (213) 955-0440 67-3510	
	(2) INFORMATION FOR SEQ ID NO:	1:	
	(B) TYPE: (C) STRANDEDNESS: (D) TOPOLOGY:	ICS: 544 nucleic acid single linear PTION : SEQ ID NO: 1:	
	GAAGGCATT GCTACAAGGT CTTCAAAC	AA TCTAAGACCT GGACAGATGC 50	0
0	AGAGAGCTTC TGCACGAAGC AGGTGAAC	GG GGGCATCTG GTCTCTATCG 10	00
	AAAGCTCCGG AGAAGCAGAC TTTGTGGG	CC AGTTGATTGC TCAGAAGATA 15	50
fin fin	AAGTCAGCCA AAATCCATGT CTGGATCG	GA CTGAGGGCTC AAAACAAAGA 20	00
Janes Brans	AAAGCAATGC AGCATAGAGT GGAGCGAT	GG CTCCAGCATC AGTTATGAGA 25	50
	ATTGGATTGA AGAAGAATCC AAAAAGTG	TC TTGGGGTGCA CATAGAGACA 30	00
25	GGGTTTCATA AGTGGGAGAA TTTTTACT	GT GAACAACAAG ATCCTTTGT 3	5(
	CTGCGAGGCA TAGTCTGAAG ATCCAGCT	GA TTGAAGTCTG GAGAAGCAAG 40	00
	GAAGCCCCCC ACCCCATCCC CCAACCCT	GC CTAGCCACAA TCTCTGCTAT 4	5(
	GCACCCTTTG CTCAACGGAT GCTCTCTC	TA GCTGGATCTG GTGTTGCTGC 5	0(
	TCCTGATGGG CCGGAAGTCA ATAAATTC	CTG CCTAGCCTGA AAAA 54	44
30	(a) TWDODWATON TOD GEO TO WO		
	(2) INFORMATION FOR SEQ ID NO	: 2:	

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:

(C) REFERENCE/DOCKET NUMBER: 233/298

129

(B) TYPE:

amino acid

(C) STRANDEDNESS:

single

(D) TOPOLOGY:

linear

5

10

15

20¹

===

îU

253

35

40

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION : SEQ ID NO: 2

Asp Cys Ser Ser Asp Trp Ser Ser Tyr Glu Gly His Cys Tyr Lys

Val Phe Lys Gln Ser Lys Thr Trp Thr Asp Ala Glu Ser Phe Cys

Thr Lys Gln Val Asn Gly Gly His Leu Val Ser Ile Glu Ser Ser

Gly Glu Ala Asp Phe Val Gly Gln Leu Ile Ala Gln Lys Ile Lys

Ser Ala Lys Ile His Val Trp Ile Gly Leu Arg Ala Gln Asn Lys

Glu Lys Gln Cys Ser Ile Glu Trp Ser Asp Gly Ser Ser Ile Ser

Lys Glu Asn Trp Ile Glu Glu Glu Ser Lys Lys Cys Leu Gly Val

His Ile Glu Thr Gly Phe His Lys Trp Glu Asn Phe Tyr Cys Glu

Gln Gln Asp Pro Phe Val Cys Glu Ala

(2) INFORMATION FOR SEQ ID NO: 3:

30 (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:

29

(B) TYPE:

amino acid

(C) STRANDEDNESS:

single

(D) TOPOLOGY:

linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION : SEQ ID NO: 3

Asp Cys Ser Ser Asp Trp Ser Ser Tyr Glu Gly His Cys Tyr Lys

- (2) INFORMATION FOR SEQ ID NO: 4:
 - (i) SEQUENCE CHARACTERISTICS: (A) LENGTH:

(B) TYPE:

amino acid

(C) STRANDEDNESS:

single

(D) TOPOLOGY:

linear

- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION : SEQ ID NO: 4

Asp Cys Pro Ser Glu Trp Ser Ser Tyr Glu Gly Phe Cys Tyr Lys Pro Phe

5

10